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RAW SECUENCE LISTING PATENT APPLICATION: US/10/073,644C DATE: 11/13/2002 TIME: 14:09:54

Input Set : A:\SEOLISTCORRECTED2.TXT Output Set: N:\CRF4\11132002\J073644C.raw

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3 <110> APPLICANT: Hudson, Debra
             van de Winkel, Jan
      5
              van Dijk, Marc
     7 <120> TITLE OF INVENTION: HUMAN MONOCLONAL ANTIBODIES TO FC ALPHA
      Я
             RECEPTOR (CD89)
     10 <130> FILE REFERENCE: MXI-211
C--> 12 <140> CURRENT APPLICATION NUMBER: US/10/073,644C
C--> 12 <141> CURRENT FILING DATE: 2002-02-11
     12 <150> PRIOR APPLICATION NUMBER: US 60/338,956
     13 <151> PRIOR FILING DATE: 2001-11-05
     15 <150> PRIOR APPLICATION NUMBER: US 60/268,075
     16 <151> PRIOR FILING DATE: 2001-02-12
    18 <160> NUMBER OF SEQ ID NOS: 8
    20 <170> SOFTWARE: FastSEQ for Windows Version 4.0
    22 <210> SEQ ID NO: 1
    23 <211> LENGTH: 357
    24 <212> TYPE: DNA
    25 <213> ORGANISM: Homo sapiens
    27 <400> SEQUENCE: 1
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    29 teetgtgeag cetetggatt cacetteagt agttatgtte tgeactgggt eegecagget 120
    30 ccaggcaagg ggctggattg ggtggcagtg atatcagatg atggaaggaa taaatacttc 180
    31 gcagactecg tgaagggeeg atteaceate tecagagaca attecaagaa caegetgtat 240
     32 ctgcaaatga acagcctgag agctgaggac acggctgtgt attactgtgt gagagaaggg 300
    33 tatageggea getggtttga etaetgggge eagggaacce tggteacegt eteetca
    35 <210> SEO ID NO: 2
     36 <211> LENGTH: 119
    37 <212> TYPE: PRT
     38 <213> ORGANISM: Homo sapiens
    40 <400> SEQUENCE: 2
    41 Gln Val Gln Leu Val Glu Ser Gly Gly Gly Val Val Gln Pro Gly Arg
    42 1
                                           10
    43 Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Ser Tyr
                   20
    44
                                        25
    45 Val Leu His Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Asp Trp Val
    46
               35
                                   40
    47 Ala Val Ile Ser Asp Asp Gly Arg Asn Lys Tyr Phe Ala Asp Ser Val
    49 Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
                           70
                                                75
    51 Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
                        85
                                           90
    53 Val Arg Glu Gly Tyr Ser Gly Ser Trp Phe Asp Tyr Trp Gly Gln Gly
```

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100
                                  105
                                                      110
55 Thr Leu Val Thr Val Ser Ser
56
          115
59 <210> SEO ID NO: 3
60 <211> LENGTH: 321
61 <212> TYPE: DNA
62 <213> ORGANISM: Homo sapiens
64 <400> SEQUENCE: 3
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66 atcacttgcc gggcaagtca gggcattagc agtgctttag cctgqtatca qcaqaaacca 120
67 gggaaagete etaageteet gatetatggt geeteeagtt tggaaggtgg ggteeeatea 180
68 aggttcagcg gcagtggatc tgggacagat ttcactctca ccatcagcag cctgcagcct 240
69 gaagattttg caacttatta ctgtcaacag tttaatagtt acccattcac tttcggccct 300
70 gggaccaaag tggatatcaa a
                                                                     321
72 <210> SEO ID NO: 4
73 <211> LENGTH: 107
74 <212> TYPE: PRT
75 <213> ORGANISM: Homo sapiens
77 <400> SEQUENCE: 4
78 Ala Ile Gln Leu Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val Glv
79 1
                   5
                                       10
80 Asp Arg Val Thr Ile Thr Cys Arg Ala Ser Gln Gly Ile Ser Ser Ala
81
              20
                                   25
82 Leu Ala Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Leu Leu Ile
83 35
                               40
84 Tyr Gly Ala Ser Ser Leu Glu Gly Gly Val Pro Ser Arg Phe Ser Gly
86 Ser Glv Ser Glv Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Gln Pro
88 Glu Asp Phe Ala Thr Tyr Tyr Cys Gln Gln Phe Asn Ser Tyr Pro Phe
                   85
                                       90
90 Thr Phe Gly Pro Gly Thr Lys Val Asp Ile Lys
91
              100
92 <210> SEO ID NO: 5
93 <211> LENGTH: 357
94 <212> TYPE: DNA
95 <213> ORGANISM: Homo sapiens
97 <400> SEQUENCE: 5
98 caggtqcagc tqqtqqaqtc tqqqqqaqqc qtqqtccaqc ctqqqaqqtc cctqaqactc 60
99 teetgtqcaq cetetqgatt cacettcagt aqetatqeta tqcactqqqt ceqecaqqet 120
100 ccaggcaagg ggctggagtg ggtggcagtt atatcatatg atggaagaaa taaagactac 180
101 gcagactccg tgaagggccg attcaccatc tccagagaca attccaagaa cacgctgtat 240
102 etgeaaatga acageetgag agetgaggae acggetgtge attactgtge gaggettgae 300
103 tggggatatg atgcttttga tatctggggc caagggacaa tggtcaccgt ctcttca
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106 <211> LENGTH: 119
107 <212> TYPE: PRT
108 <213> ORGANISM: Homo sapiens
110 <400> SEOUENCE: 6
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111 Gln Val Gln Leu Val Glu Ser Gly Gly Val Val Gln Pro Gly Arq
112 1
                                        10
113 Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Ser Tyr
114
                20
                                    25
115 Ala Met His Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
116
           35
                                40
117 Ala Val Ile Ser Tyr Asp Gly Arg Asn Lys Asp Tyr Ala Asp Ser Val
118
                            55
119 Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
120 65
121 Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val His Tyr Cys
123 Ala Arq Leu Asp Trp Gly Tyr Asp Ala Phe Asp Ile Trp Gly Gln Gly
124
                                    105
                                                        110
125 Thr Met Val Thr Val Ser Ser
126
           115
128 <210> SEO ID NO: 7
129 <211> LENGTH: 327
130 <212> TYPE: DNA
131 <213> ORGANISM: Homo sapiens
133 <400> SEQUENCE: 7
134 gaaattgtgt tgacgcagtc tccaggcacc ctgtctttgt ctccagggga aagagccacc 60
135 eteteetgea gggccagtea gagtgttage ageagetact tagcetggta ccagcagaag 120
136 cctggccagg ctcccaggct cctcatctat ggtgcatcca gcagggccac tggcatccca 180
137 gacaggitca giggcagigg gictgggaca gacticactc tcaccatcag cagactggag 240
138 cctqaaqatt ttqcaqtqta ttactqtcaq caqtatqqta qctcacctcc qtacactttt 300
139 ggccagggga ccaagctgga gatcaaa
141 <210> SEO ID NO: 8
142 <211> LENGTH: 109
143 <212> TYPE: PRT
144 <213> ORGANISM: Homo sapiens
146 <400> SEOUENCE: 8
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148 1
149 Glu Arg Ala Thr Leu Ser Cys Arg Ala Ser Gln Ser Val Ser Ser Ser
151 Tyr Leu Ala Trp Tyr Gln Gln Lys Pro Gly Gln Ala Pro Arg Leu Leu
           35
153 Ile Tyr Gly Ala Ser Ser Arg Ala Thr Gly Ile Pro Asp Arg Phe Ser
       50
155 Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Arg Leu Glu
                       70
                                            75
157 Pro Glu Asp Phe Ala Val Tyr Tyr Cys Gln Gln Tyr Gly Ser Ser Pro
159 Pro Tyr Thr Phe Gly Gln Gly Thr Lys Leu Glu Ile Lys
```

VERIFICATION SUMMARY

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L:12 M:270 C: Current Application Number differs, Replaced Current Application No L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date